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ABSTRACT: Thermal treatment of proteinaceous foods generates heat-induced Maillard reaction substances including toxic advanced glycation end products (AGEs) and heterocyclic amines (HAs). It is known that plant phenolic compounds may influence Maillard reaction. This study investigated the impact of lingonberry leaf extracts on the formation of N^ε-(carboxymethyl)lysine (CML) and N^ε-(2-furoylmethyl)-L-lysine (furosine) in milk model system and HAs in meat-protein and meat model systems. In addition, lingonberry leaf extracts obtained by different solvents were characterized by radical scavenging, Folin-Ciocalteu assays and ultra-high pressure liquid chromatography quadruple-time-of flight mass spectrometry (UPLC-qTOF-MS). Water extract (WE) stronger suppressed CML than furosine formation in milk model system: CML levels were reduced by nearly 40%. Moreover, quinic acid and catechin which were abundant in WE, were effective in inhibiting CML and furosine formation. WE and acetone extract (AE) at 10 mg/mL significantly inhibited HAs formation in both model systems. However, higher suppressing effect on HAs formation showed AE which had lower antioxidant capacity and total phenolic content values than WE. WE contained higher amounts of hydroxycinnamic acids, proanthocyanidins and flavonols, while AE was richer in flavan-3-ols and arbutin derivatives. It indicates that the composition of phenolics might be a major factor for explaining different effect of extracts from the same plant on HAs formation. In general, the results suggest that lingonberry leaves is a promising source of phytochemicals for inhibiting toxic Maillard reaction products and enriching foods with plant bioactive compounds.

Keywords: *Vaccinium vitis-idaea* L. leaves; Maillard reaction; CML; furosine; heterocyclic amines

58 **Practical Application:**

59 The increased consumption in processed foods has been linked with the increased risks of
60 various diseases, while thermal food processing is required to develop flavour, insure safety
61 and extend shelf life. Therefore, developing effective technological means for inhibiting the
62 formation of heat-induced toxic substances is an important task. This study showed a potential
63 of lingonberry leaf extracts containing health beneficial phytochemicals to suppress the
64 formation of toxic Maillard reaction products during heating of milk and meat.

Introduction

Thermal treatment of raw milk or meat during processing is an essential step in reducing microbiological contamination causing foodborne diseases and extending product shelf life. On the other hand, thermal treatment results in the formation of undesirable compounds demonstrating toxicity and allergenicity. The majority of those modifications are linked to the Maillard reaction occurring between carbonyl groups of reducing sugars and free amino groups of amino acids (Nursten, 2005).

Thermal treatment of raw milk in such processing steps as separation, normalization, homogenization, pasteurization, sterilization or ultra-heat treatment induces Maillard reaction mainly between lactose and lysine residues and generates the advanced glycation end products (AGEs). The formation of AGEs increases with increasing heating time and temperature. The presence of AGEs in foods contributes to the increased inflammation and oxidative stress (Uribarri et al., 2007), which are linked to the recent epidemics of diabetes and cardiovascular diseases (Birlouez-Aragon et al., 2010; Sandu et al., 2005). A range of dietary AGEs compounds have been reported, while the present study is focused on N^ε-(carboxymethyl)lysine (CML) and indirect marker, N^ε-(2-furoylmethyl)-L-lysine (furosine). CML is one of the most studied AGE, which is present in both biological systems, such as plasma, urine, tissues, skin collagen and in many heat-processed foods (Nguyen, van der Fels-Klerx, & van Boekel, 2014). It is generated from the oxidation of Amadori product (AP) or direct reaction of lysine ε-amino group with glyoxal produced by degradation/oxidation of sugars during thermal treatment and lipid peroxidation. Furosine is an indirect marker of AGEs indicating early stage of Maillard reaction

and is formed from the acid hydrolysis of the APs (e.g. fructosyl-lysine) (Henle, Zehetner, & Klostermeyer, 1995).

Thermal treatment of meat, particularly by deep-frying, oven-frying, roasting, grilling and searing generates the formation of heterocyclic amines (HAs) possessing mutagenic and carcinogenic properties. They are formed mainly as Maillard reaction products from amino acids, creatin(in)e and reducing sugars (Skog, Johansson, & Jägerstad, 1998). Depending on the cooking method, heating temperature and time, as well as on the content of precursors the formation of different HAs are favoured during thermal treatment (Gibis & Weiss, 2015; Linghu, Karim, & Smith, 2017; Yu et al., 2018). 2-Amino-1-methyl-6-phenylimidazo[4,5-b] pyridine (PhIP) is the most abundant mutagenic HA of meat, while the chicken was reported as the most susceptible type of meat for the formation of this HA (Gibis & Weiss, 2015). It was observed that PhIP is carcinogenic in the rat colon, mammary gland, prostate, and breast (Shirai et al., 1997). It was also shown that many HAs possess carcinogenic/mutagenic/genotoxic effects both *in vitro* (Dumont et al., 2010) and *in vivo* (Chao et al., 2005; Shirai et al., 1997). Recent studies have supported positive associations of HA's with colorectal adenoma risk (Budhathoki et al., 2015; Khan et al., 2019) and increased oxidative stress in humans (Carvalho et al., 2015).

Phenolic compounds have been shown to possess inhibiting effects on the formation of AGEs and HAs (Puangsombat, Jirapakkul, & Smith, 2011). These effects have been explained by different reaction mechanisms, namely radical scavenging (Kikugawa, 1999), amine group blocking (Guerra & Yaylayan, 2014), scavenging dicarbonyl intermediates (Totlani & Peterson, 2005) or reactive carbonyls produced from amino acid degradation (Cheng et al., 2009) and lipid oxidation (Hidalgo, Delgado, & Zamora, 2017). For instance, catechin (Kokkinidou &

Peterson, 2014), (–)-epicatechin (Totlani & Peterson, 2006), (–)-epigallocatechin-3-gallate (EGCG) (Bin, Peterson, & Elias, 2012) inhibited AGEs by trapping dicarbonyl compounds. EGCG inhibited PhIP by scavenging reactive carbonyls from phenylacetaldehyde degradation (Cheng et al., 2009). Moreover, the structure of phenolics was explored as an important factor in the inhibition of HAs formation (Salazar, Arámbula-Villa, Hidalgo, & Zamora, 2014). Therefore, phenolic antioxidant-rich natural plant extracts may be useful additives in proteinaceous foods for inhibiting toxic Maillard reaction products.

Lingonberry (*Vaccinium vitis-idaea* L., Ericaceae) is one of the most important wild berry plant in the forests of the Nordic countries. Lingonberry leaves have been used in traditional medicine for their diuretic, astringent, and antiseptic properties to treat urinary tract infections. Lingonberry leaves contain many classes of flavonoids such as flavan-3-ols, flavanols, hydroxycinnamic acids, and proanthocyanidins (Hokkanen, Mattila, Jaakola, Pirttilä, & Tolonen, 2009; Ieri, Martini, Innocenti, & Mulinacci, 2013; Liu et al., 2014). According to European Commission Novel Food Catalogue lingonberry leaves have been authorized as food supplements. In addition, *V. vitis-idaea* berry extracts demonstrated antiglycation activity (Beaulieu et al., 2010), while there are no reports on antiglycation properties or HAs inhibition by lingonberry leaf extracts.

So far as milk and meat are very important foods for adequate human nutrition, particularly in supplying high value proteins and various bioactive micronutrients, the application of plant antioxidants for reducing the formation of undesirable toxic compounds during heat treatment might be a promising approach in achieving better equilibrium between the positive and negative nutritional aspects in the consumption of heat processed proteinaceous foods.

Therefore, this study was aimed at investigating the effect of lingonberry leaf extracts on the formation of CML and furosine in milk model system and HAs in meat-protein and meat model systems. The antioxidative potential of different lingonberry leaf extracts obtained by different solvents from deodorized and non-deodorized material was characterized by radical scavenging capacity, the total phenolic content and composition of phenolic compounds.

Materials and methods

Chemicals

2,2-Diphenyl-1-picrylhydrazyl (DPPH^{*}), 2,2-azinobis-3-ethyl benzothiazoline-6-sulphonic acid (ABTS), 6-hydroxy-2,5,7,8,-tetramethylchromane-2-carboxylic acid (Trolox), 2,2'-azobis(2-methylpropionamidine) dihydrochloride (AAPH), diethylene glycol, sodium borohydride, boric acid, nonafluoropentanoic acid (NFPA), trichloroacetic acid (TCA), D (+)-lactose monohydrate, acetonitrile (HPLC grade), rutin, quercetin-3-*O*-glucoside, quinic acid, *p*-coumaric, caffeic and gallic acids, Folin–Ciocalteu phenol reagent and Milli-Q water were from Sigma-Aldrich (St. Louis, MO, USA). (+)-Catechin reference was purchased from Chromadex (Irvine, California, USA), chlorogenic acid, L-phenylalanine and glucose were from Roth (Karlsruhe, Germany); fluorescein sodium salt (FL) from Fluka Chemicals (Steinheim, Germany). Heterocyclic amines, MeIQ (2-amino-3,4-dimethylimidazo[4,5-f]quinolone), MeIQx (2-amino-3,8-dimethylimidazo[4,5-f] quinoxaline), Trp-P-1 (3-amino-1,4-dimethyl-5H-pyrido[4,3-b]indole) and PhIP were purchased from Toronto Research Chemicals Inc. (Toronto, Canada). CML, d₂-CML and furosine were purchased from PolyPeptide Laboratories (Strasbourg, France). Creatinine was from Alfa Aesar (Karlsruhe, Germany). Bond Elut PRS LRC cartridges and Bond-

Elut C18 cartridges were purchased from Agilent Technologies (Lake Forest, CA, USA). Chromabond XTR cartridges were from Macherey-Nagel GmbH & Co. (Bethlehem, PA, USA). Oasis HLB cartridges for CML and furosine stable isotope dilution assay were purchased from Waters Corporation (Milford, CA, USA).

Plant material and preparation of extracts

Vaccinium vitis-idaea L. leaves were collected in the forest of Mažeikiai (Telšiai district, Lithuania) from June to mid-September. The plants were dried at 40 °C and ground in an ultra-centrifugal rotor mill Retsch ZM200 (Retsch GmbH, Haan, Germany) using 0.5 mm sieve. Powdered lingonberry leaves (200 g) were placed in a 3 L round-bottom flask, connected with Clevenger-type apparatus, diluted with 1.5 L of water and hydrodistilled during 3 h for removing volatile compounds. Then, water phase was separated from the solids by filtering through the cotton and freeze-dried. The product obtained is indicated as water extract (WE). The solid residues, which are called deodorized lingonberry leaves, were dried at 40 °C in a drying oven. Then, raw and deodorized lingonberry leaves were subjected to pressurised liquid extraction with methanol and acetone in a Dionex ASE 350 apparatus (Sunnyvale, CA, USA) at the following parameters: 70 °C, 10 MPa, 3 cycles, 5 min each, 100% flush volume, purging with nitrogen. Organic solvents were evaporated at 40 °C in a rotary vacuum evaporator (Büchi, Flawil, Switzerland). Acetone, as an aprotic lower polarity solvent, and methanol, as a protic high polarity solvent, were selected in order to evaluate what kind of compounds may be recovered by using these solvents and what extracts (containing lower or higher polarity phytochemicals) possess the strongest radical scavenging properties. In total, five extracts were

produced: three extracts from the raw (WE, ME, AE) and two extracts from the deodorized lingonberry leaves (DME and DAE).

Evaluation of antioxidant potential of lingonberry leaves

Folin-Ciocalteu assay

Total phenolic content was determined using Folin-Ciocalteu method (Singleton, Orthofer, & Lamuela-Raventós, 1999). Extract solution (0.2 mL) was added to 1.0 mL of Folin–Ciocalteu reagent (1:10, v/v) and was shaken for 5 min; then 0.8 mL of 7.5 mg/g sodium carbonate solution was added and after incubation in the dark for 1.5 h, the absorbance was read at 760 nm in a HALO RB-100 UV–Vis spectrophotometer (Dynamica, Switzerland). The results were expressed as mg gallic acid equivalents (GAE) in g of extract dry weight (edw) and in plant dry weight (pdw).

DPPH• scavenging assay

DPPH• scavenging was determined according to Brand-Williams, Cuvelier, & Berset (1995). Briefly, the decrease in absorbance was recorded at 517 nm in a FLUOstar Omega microplate reader (BMG Labtech, Durham, NC) over 35 min after the addition of 7.5 µL of extract or trolox solution in 96-well microplates with 300 µL of methanolic DPPH• solution (60 µM). The results were expressed as effective concentrations EC₅₀ (mg/mL) required to decrease the initial DPPH• concentration in the reaction mixture by 50%.

ABTS^{•+} scavenging assay

The ABTS^{•+} scavenging (decolouration) capacity was determined according to Re et al. (1999). Briefly, after adjusting the absorbance of ABTS^{•+} solution to 0.70 ± 0.02 at 734 nm with phosphate buffer (pH 7.4), the measurements were performed in microplate reader after mixing 3 μ L of extract or trolox solutions with 297 μ L of diluted ABTS^{•+} solution. Trolox equivalent antioxidant capacity (TEAC) was used to express ABTS^{•+} scavenging capacity in mM TE/g edw and pdw.

Oxygen radical absorbance capacity (ORAC) assay

The ORAC assay was performed as described by Huang, Ou, Hampsch-Woodill, Flanagan, & Prior (2002). Briefly, 25 μ L of extract or trolox solution were mixed with 150 μ L of 96 mM FL in microplate, which further was incubated at 37 °C for 15 min. Then, 26 μ L of 240 mM AAPH was added and microplate was shaken for 30 s. The fluorescence was recorded ($\lambda_{\text{ex}} = 493$ nm, $\lambda_{\text{em}} = 515$ nm) every 66 s during 80 min. The ORAC values are expressed as mM TE/g edw and pdw.

Analysis of phenolic compounds

Phenolic compounds were analyzed by ultra-high pressure liquid chromatography with quadrupole-time of flight mass spectrometry (UPLC-qTOF-MS) using an Acquity UPLC system (Waters Corporation) and Brüker maXis UHR-TOF mass spectrometer (Brüker Daltonics, Bremen, Germany). Separation was performed using 1.7 μ m Acquity BEH-C18 (50 \times 2.1 mm) column at a flow rate of 0.4 mL/min (injection volume 2 μ L). The eluents were 0.1% formic acid (A) and acetonitrile (B) at the following gradient: 0–2 min, 10–22% B, 2–7 min, 22%–50% B, 7–10 min, 10% B. Mass spectra were acquired in a negative electrospray ionization mode (ESI) by

full scan acquisition covering a range of m/z 100-800 at the following parameters: capillary voltage, 4000 V; collision cell energy, 8 eV, dry gas temperature, 200 °C with a flow rate of 10 L/min; nebulizer pressure, 2 bar; drying and nebulizing gas, nitrogen. Peak identification was performed by comparing retention times (t_R) with those of standards or and accurate m/z values from which the molecular ion formulas were calculated for each compound. Mass fragmentation was performed with nitrogen as a collision gas at 40–45 eV collision energy.

Determination of effects of lingonberry leaf extracts on Maillard reaction products

Preparation of milk model system and determination of furosine and CML

Milk model system was prepared by dissolving (w/w) 1.2% skimmed milk powder, 2.5% whey protein (both from Lactalis Ingredients, Bourgbarré, France) and 5% lactose monohydrate in 87.9% of water, and then homogenizing (60 °C, 16 MPa, 3 passes) with 3.3% of palm oil (Kerfoot Groop, Yorkshire, UK) and 0.1% of soy lecithin Solec (DuPont, Dangé Saint Romain, France). WE, quinic acid and catechin were added at 0.05, 0.1 and 0.3 mg/mL concentrations, while the controls (also in other model systems) were without any additives. The samples were heated at 140 °C for 0, 30 and 45 s. Furosine and CML were extracted and determined as described by Delatour et al. (2009) and Fenaille et al. (2006) with slight modifications reported by Račkauskienė et al. (2015).

Preparation of meat-protein model system and determination of HAs

Meat-protein model system consisting of 0.4 mM phenylalanine, 0.4 mM creatinine and 0.2 mM glucose in 3 mL of diethylene glycol (14% of water) was used as described by Wong, Cheng,

& Wang (2012). WE and AE powders were added to the mixture at 0.1, 1, 5, and 10 mg/mL levels. The samples were heated at 128 °C in the closed stainless steel vials (homemade from stainless steel type EN 1.4401) for 120 min, afterwards cooled in an ice-water bath for 10 min and stored at –18 °C. After extraction of HAs (Wong, Cheng, & Wang, 2012) the eluate was dried under nitrogen and re-dissolved in 400 µL of methanol before analysis.

PhIP was determined using the same UPLC system as for phenolic compounds. Separation of HAs was performed using Acquity BEC-C18 column thermostated at 40 °C. The mobile phase was composed of 30 mM formic acid/ammonium formate (pH 4.75) in deionized water (A) and acetonitrile (B) delivered at 0.4 mL/min (injection volume 5 µL) at the following gradient: 0–0.15 min, 5% B; 0.15–2.5 min, 5–30% B; 2.5–3.0 min, 30–60% B; 3–5 min, 60% B and 5–7 min 60–100% B. Positive ESI was used for mass analysis and MS parameters were the same as described above. The full scan MS data were recorded in the range of 100 to 800 m/z. PhIP was quantified using calibration curve prepared from its solutions in methanol (1 - 800 ng/mL) and following the whole procedure described above. The limits of detection (LOD) and quantification (LOQ) were 1 and 3 ng/mL, respectively, while average PhIP recovery was approx. 61%.

Preparation of meat model system and determination of HAs

A meat model system consisting of 0.1 g of homogenized freeze-dried beef meat purchased in the local market (Kaunas, Lithuania) and 1 mL of diethylene glycol (14% of water) was used (Messner & Murkovic, 2004). WE and AE powders were added to the mixture at 0.1, 1, 5, and 10 mg/mL concentrations. The samples were placed in the stainless steel vessels, which were

closed and heated at 180 °C for 30 min. After heating the vessels were cooled on ice (10 min) to terminate the reaction and the samples were stored at –18 °C prior to extraction. The samples were then transferred to 50 mL glass beakers with 15 mL of 1 M NaOH and homogenized by magnetic stirring for 60 min at 500 rpm. Next, 5 mL of alkaline solution were mixed with diatomaceous earth in Chromabond XTR cartridges, and extracted (Wong et al., 2012). The eluate was dried under nitrogen and re-dissolved in 100 µL of methanol before analysis.

HAs were analysed on a Waters Acquity UPLC H-Class system equipped with triple quadrupole spectrometer Waters Xevo TQ-S in a positive ESI mode. The compounds were separated in an Acquity BEH-C18 column at a flow rate of 0.8 mL/min (injection volume 2 µL). The mobile phase was 30 mM formic acid-ammonium formate buffer (A) and acetonitrile (B) eluted at the following gradient: 0–0.1 min, 5% B; 0.1–1.5 min, 5–30% B; 1.5–1.8 min, 30–60% B; 1.8–1.85 min, 60% B; 1.85–2.4 min, 60–95% B; 2.4–2.9 min, 5% B. The quantitation was carried out in multiple reaction monitoring (MRM) mode using transitions as follows: 225 → 210 for PhIP; 213 → 198 for MeIQ; 214 → 199 for MeIQx; 212 → 195 for TrP-P-1. MRM conditions were automatically optimized with 100 ng/mL HAs using the Intellistart function system. MS/MS parameters were as follows: cone voltage, 50 V; capillary voltage, 3.0 kV; desolvation temperature, 350 °C; cone gas flow rate, 150 L/h (Nitrogen); desolvation gas flow rate, 650 L/h (Nitrogen); collision gas flow rate, 0.13 mL/min (Argon); collision energy, 35 eV. Data acquisition was carried out by MassLynx 4.1 software. Quantitative analysis was performed using Waters TargetLynx™ software. Calibration curves ranged from 1 to 500 ng/mL. The LOD and LOQ for standard solutions were calculated based on signal-to-noise ratios of 3:1 and 10:1,

respectively. The LOD and LOQ values (ng/mL) were 1 and 2 for PhIP, 2 and 5 for MeIQ, 2 and 5 for MeIQx, and 2 and 6 for Trp-P-1, respectively. The recovery ranged from 67 to 86%.

Statistical analysis

Antioxidant characteristics are reported as means \pm standard deviations (SD) from 3 replicate measurements. Milk, meat-protein and meat model systems were prepared in duplicate, while extraction procedure was performed in duplicate for each replicate. Each replicate was analysed in triplicate in LC-MS/MS and UPLC systems. Multivariate analysis with Turkey test as a post hoc analysis was used for statistical data assessment ($P < 0.05$).

Results and discussion

Evaluation of antioxidant properties of lingonberry leaves

Extraction yields and total phenolic content of lingonberry leaf extracts

In case of raw lingonberry leaves the highest and the lowest extraction yields were obtained with methanol and acetone, respectively, whereas deodorized lingonberry leaves gave 2-fold lower extract yields than raw lingonberry leaves (Table 1). In general, the yields decreased in the following order: ME > WE > AE > DME > DAE. The differences in the ME and WE yields were not large; both solvents are high polarity compounds, whereas methanol is known as one of the most effective solvents for the extraction of low sugar content plant materials. Moreover, methanol extraction was performed at the increased pressure, which facilitates diffusion and solubilization processes. Acetone is less effective solvent for higher polarity constituents, while DME and DAE were prepared from the solid hydrodistillation residue, which has already been

extracted by water. The highest amounts of total phenolics expressed in mg GAE/g pdw were recovered with ME followed by WE, AE, DME, and DAE. More than 2-fold lower total phenolic content was determined for the deodorized than for the raw lingonberry leaf extracts. Recent study (Bujor, Ginies, Popa, & Dufour, 2018) reported total phenolic content for lingonberry leaves in the range of 85.3 - 114.6 mg GAE/g DM, which is in agreement with our data (57.42±1.19–91.15±1.3 mg GAE/g pdw).

Radical scavenging properties of lingonberry leaf extracts

DPPH[•], ABTS^{•+} scavenging and ORAC values are presented in Table 1. DAE was the weakest DPPH[•] scavenger while the differences between EC₅₀ values of WE, ME, AE, and DME were not significant in this assay. It was also observed that lingonberry leaf extracts scavenged DPPH[•] rather slowly; the reaction between antioxidants and radical was completed in approx. 35 min. In ABTS^{•+} and ORAC assays the antioxidant capacity of extracts was decreasing in the following order: WE > ME > AE > DME > DAE. A significant correlation exists between total phenolic content (mg GAE/g pdw) and DPPH[•] scavenging ($r = -0.70$), ABTS^{•+} scavenging ($r = 0.87$) and ORAC ($r = 0.95$) values.

In general, high polarity protic solvents methanol and water gave higher yields, total phenolic content and antioxidant capacity values than lower polarity solvent acetone. It indicates that lingonberry leaves contain higher amounts of polar polyphenolic compounds. Therefore, extraction yields, total phenolic content and antioxidant capacity values of extracts obtained from the deodorized material were remarkably lower in comparison to the extracts from the raw material; the main part of water-soluble compounds remains in WE after hydro-

distillation of lingonberry leaves. It should be noted that comprehensive *in vitro* evaluation of radical scavenging capacity of different lingonberry leaf extracts is reported for the first time.

Comparison of phenolic profiles of different lingonberry leaf extracts

The UV chromatograms and the relative amounts (based on UPLC-qTOF-MS peak areas) of phenolic compounds in different lingonberry leaf extracts are presented in Figure 1 and Table 2. Twenty phenolic compounds were positively or tentatively identified in lingonberry leaf extract. Authentic standards were used for identification of catechin, quinic, chlorogenic, caffeic, citric and *p*-coumaric acids, rutin and quercetin-3-O-glucoside. Other compounds were tentatively identified by comparing the exact masses, fragmentation patterns, t_R by using databases such as METLIN and Human Metabolome (HMDB) (Benton, Wong, Trauger, & Siuzdak, 2008) and previously reported data for those compounds in lingonberry leaves (Ek, Kartimo, Mattila, & Tolonen, 2006; Liu et al., 2014). It should be noted that so far as standards were not available for these compounds their identification is assumed as tentative. ME and AE demonstrated lower recovery of phenolics, expressed in $AU \times 10^{-7}/mg$ edw, than WE by 37 and 51%, respectively. Lower recovery of phenolics by acetone may be explained by the prevalence of higher polarity compounds in lingonberry leaves, which are better soluble in protic solvents. Finally, DME and DAE contained lower amounts of total phenols by 53 and 81% than WE, respectively. The content of phenolics in deodorized plant extracts was considerably lower comparing with the extracts from the raw lingonberry leaves, because the main part of polar compounds was dissolved in water during hydro-distillation. It indicates that after water

extraction of raw lingonberry leaves some part of extractable phenolics still remained in deodorized lingonberry leaves and were recovered by methanol and acetone.

The main classes of phenolics in WE were hydroxycinnamic acids and flavonols while arbutin derivatives and flavan-3-ols were more abundant in ME and AE. The dominant flavonols in all extracts were pentosides and hexosides of quercetin, which is in agreement with previous studies (Ek et al., 2006, Liu et al., 2014). Exact mass and fragmentation patterns indicated that the compounds **16** and **17** may be quercetin-3-*O*-xyloside and quercetin-3-*O*-arabinoside. This assumption is also supported by the order of their elution, which was determined in previous study (Liu et al., 2014). Caffeic acid **7**, *p*-coumaric acid **8** and quercetin-pentoside **16** were found only in WE, DME and DAE indicating that detectable amounts of these compounds were released after hydro-distillation. Also, citric acid **4**, chlorogenic acid **6**, *B*-type proanthocyanidin **12** and rutin **14** were found only in WE.

The differences in the yields, antioxidant capacity and total phenols for WE and ME were not considerable. This is a positive result indicating that green and cheap solvent water may be used for preparing phenolics-rich extracts from lingonberry leaves on industrial scale. However, the profile of phenols in WE was different compared to ME and AE, while in the latter extracts it was quite similar. WE and AE were chosen for further evaluation of their possible inhibition of toxic Maillard reaction products.

Effects of lingonberry leaves on Maillard reaction products

The effect of WE, quinic acid and catechin on furosine and CML formation

In this experiment, the antiglycation activity of WE, quinic acid and catechin was accessed through the furosine and CML determination in milk model system. Since WE exhibited the highest antioxidant potential it was chosen for this experiment. To identify possible active components in the lingonberry leaves, the standards of quinic acid and catechin, were further evaluated. Quinic acid was among the major constituents in WE, while catechin was the strongest radical scavenger as it was determined by the on-line HPLC-DPPH[•] scavenging method (data not shown).

The effect of WE on furosine and CML formation is represented in Figure 2A1 and A2, respectively; the former was better inhibited at lower, while the latter one at higher concentrations of WE. For instance, at 0.05 and 0.1 mg/mL WE reduced furosine level by 32 and 21% after 30 s of heating and by 18 and 27% after 45 s of heating, respectively. CML formation was reduced by 38 and 19% after 30 s of heating and by 42 and 38% after 45 s of heating at 0.1 and 0.3 mg/mL WE addition, respectively.

The effect of quinic acid and catechin on furosine and CML formation is presented in Figure 2B and C, respectively. Quinic acid and catechin reduced the levels of furosine in milk model system after 30 s heating by 35% at all concentrations; while after 45 s heating, its level was reduced by approx. 10 and 40% at 0.05 mg/mL and 0.3 mg/mL concentrations, respectively. The formation of CML was even more efficiently inhibited than furosine. The average decrease of CML at both heating times was approx. by 31, 29, and 43%, and 31, 39, and 51% when 0.05, 0.1 and 0.3 mg/mL of quinic acid (Figure 2B2) and catechin (Figure 2C2) were added, respectively. The reduction of CML and furosine by catechin can be explained through two pathways: first, through the reaction between amino groups and *o*-quinoidal moieties forming via B ring

oxidation; and second, through the trapping of carbonyl-containing sugar fragments at highly activated A ring. There are no reports showing suppressing effect of quinic acid on Maillard reaction products formation in milk. However, the protecting effect against protein carbonylation was reported previously (Yoshimura et al., 2016). Quinic acid is a cyclic hydroxyl-acid that is present in various fruits and vegetables; therefore, it would be interesting to continue the studies for clarifying reducing effect of quinic acid on the formation of Maillard reaction products.

In general, the inhibition effect of WE on Maillard reaction products formation depended on heating time and extract concentration. The reduction of both Maillard reaction products can be associated with the above-described ways. Given their antiglycation activity and relatively high abundance in leaves, catechin and quinic acid may be suggested as the main active compounds responsible for CML and furosine inhibition. Catechin, quercetin-3-*O*-galactoside and cyanidin-3-*O*-glucoside were reported as the main contributors to the antiglycation properties of *V. vitis-idaea* berry extracts (Beaulieu et al., 2010).

The effect of WE and AE on the formation of HAs

In this experiment, the effects of WE and AE of lingonberry leaves on the formation on HAs were tested in two model systems: meat-protein and meat. In a meat-protein model system, the interference of meat matrix components (lipids, proteins, water) was eliminated, by using only the substances, which are necessary for the formation of HAs. As only phenylalanine was used, mainly PhIP formation was generated; therefore the effect of lingonberry leaf extracts on the formation of PhIP was evaluated in meat-protein model system. Meat model system

consisted of a real meat-like matrix, from which only the water factor was eliminated; in this case the formation of HAs was evaluated measuring several typical to roasted beef HAs.

The effect of lingonberry leaves extracts on PhIP formation in meat-protein model system is shown in Figure 3. The amounts of PhIP significantly increased when WE were added in the range of 0.1-5 mg/mL and decreased by 20% with the addition of 10 mg/mL; while significant effect of AE was observed only at 10 mg/mL when PhIP level was reduced by 40%. Better inhibitory effect observed in case of AE, possessing weaker radical scavenging and total phenolic content than WE, may seem somewhat unexpected. According to our results and previously reported observations (Hidalgo & Zamora, 2018), it may be assumed that different effect of WE and AE on PhIP formation could be due to the different composition of phenolics: WE contained more hydroxycinnamic acids, proanthocyanidins and flavonols than AE. The amounts of hydroxycinnamic acids as quinic, protocatechuic, chlorogenic, caffeic and *p*-coumaric were in traces or even undetectable in AE. Phenolic acids, e.g. chlorogenic (Cheng, Chen, & Wang, 2007), were shown to be ineffective inhibitors or even enhancers of PhIP formation. Moreover, AE comparing to WE contained higher amounts of flavan-3-ols such as catechin and chinchonain 1 and arbutin derivative (2-O-caffeoylarbutin). Green tea catechins in many studies were reported as effective inhibitors of HAs (Cheng et al., 2009; Quelhas et al., 2010). Those differences in the composition of phenolics explain different effect of lingonberry leaf extracts on PhIP formation in the analysed meat-protein model system.

The effects of lingonberry leaves extracts on the formation of four HAs in meat model system are presented in Figure 4. In this model four HAs were quantified; the dominant compound was PhIP (266.84±6.67 ng/g of beef dry weight), whereas MeIQ, MeIQx and Trp-P-1

were found in remarkably lower amounts. Trp-P-1 is a carboline type HA and in cooked meat usually forms at higher temperatures ($> 200\text{ }^{\circ}\text{C}$); however, probably due to prolonged heating (120 min) it was also formed in the used beef-model system. WE and AE reduced PhIP concentration dose-dependent by 3.24, 10.55, 39.29 and 14.42, 26.85, 37.43%, respectively. Consequently, the opposite effects of WE addition on the formation of PhIP were observed in the more real-like meat model system: WE did not promote PhIP formation. It may be explained by the different formation mechanisms of PhIP in chemical and real-like meat model systems, as was reported previously (Cheng et al., 2007). However, the levels of MeIQ, MeIQx and Trp-1 were increased in the samples with lingonberry leaf extracts and WE showed stronger promoting effect than AE. The addition of 10 mg/mL of WE showed the highest promoting effect on the formation of MeIQ, MeIQx and Trp-1 and the highest reducing effect on PhIP formation. These results might be explained by different HAs formation mechanisms, which largely depend on their structures. For instance, MeIQ belongs to imidazoquinoline, while MeIQx to imidazoquinoxaline class of Has. The precursors of MeIQ and MeIQx are pyridine and pyrazine free radicals, respectively, forming *via* Maillard reaction between sugars and amino acids during Strecker degradation. Catechin and EGCG influenced the formation of pyrazinium radicals by reacting with imine intermediates: moreover, at low concentrations (5-20 mM) it increased and at high concentrations ($\geq 50\text{ mM}$) decreased their formation. Thus, the formation of MeIQ and MeIQx could be enhanced through the generation of pyridine and pyrazine radicals. As PhIP belongs to imidazopyridine type compound, it is formed from the condensation of phenylacetaldehyde and creatinine. EGCG has been shown as an effective

inhibitor of phenylacetaldehyde, which is an important intermediate product in PhIP formation (Cheng et al., 2009).

Based on the obtained data, we can assume that WE may act as a generator of Trp-P-1, MelQ and MelQx precursors, and as a scavenger of intermediates in the formation of PhIP in meat model system. In general, AE was more potential additive for reducing HAs formation in both model systems.

Conclusions

The results of our study indicate that lingonberry leaf extracts possess the ability of tuning Maillard reaction in food models. WE inhibited CML and furosine formation in milk model system during its heating; moreover the inhibitory effect was stronger for the formation of CML than furosine. This finding suggests that WE is stronger inhibitor of the advanced stage Maillard reaction products compared to the early stage products. Phenolic compounds such as catechin and quinic acid, which were abundant in WE, showed strong suppressing effect on CML and furosine formation in milk model system; consequently, these compounds may be important lingonberry leaves extract constituents responsible for CML and furosine inhibition. Results of extract's effects in meat models systems suggest that inhibition effect is more related to the phenolic compounds profile than to the radical scavenging capacity or total phenolic content. Accordingly, it may be preliminary hypothesized that hydroxycinnamic acids might act as enhancers, while flavan-3-ols and arbutin derivatives as inhibitors of HAs. Moreover, effective phytochemical composition of such extracts could be purposively designed by using different organic solvents and procedures for extractions. In general, the results obtained show that

lingonberry leaf extracts are promising ingredients for their applications in foods both for controlling the formation of Maillard reaction products during thermal processing and for enriching them with natural antioxidants possessing health benefits.

In addition, the results supports the need of further studies for the evaluation of the mechanisms of the involvement of plant phytochemicals in the Maillard reaction to ensure optimal application of natural plant origin ingredients in suppressing the formation of heat-induced toxic compounds in foods.

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Author Contributions

I. Račkauskienė performed the major part of experimental work (data acquisition, statistical analysis and data interpretation) and prepared the manuscript draft and finalized it after approval by the other co-authors. P.R. Venskutonis conceptualized and designed the study, critically reviewed the data and finally approved the manuscript. A. Pukalskas, A. Fiore and A.D. designed the study, performed part of experimental work, interpreted data, critically reviewed it and finally approved the manuscript.

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650 **Table 1–The yield, total phenolic content and antioxidant activity of different extracts of lingonberry leaves.**

LL	Yield	TPC	TPC	DPPH, EC ₅₀	ABTS	ABTS	ORAC	ORAC
Extract	(g/100 g pdw)	(mg GAE/edw)	(mg GAE/g pdw)	(mg/mL)	(mM TE/g edw)	(mM TE/g pdw)	(mM TE/g edw)	(mM TE/g pdw)
WE	32.96±1.90 ^d	252.22±4.40 ^c	83.12±1.45 ^d	2.40±0.01 ^a	11.16±0.55 ^d	3.68±0.18 ^d	10.63±0.88 ^e	3.50±0.29 ^e
ME	36.38±2.51 ^e	250.56±3.62 ^c	91.15±1.31 ^e	2.36±0.00 ^a	6.05±1.55 ^c	2.20±0.57 ^c	7.44±1.05 ^d	2.71±0.38 ^d
AE	23.69±2.60 ^c	242.42±5.05 ^c	57.42±1.19 ^c	2.52±0.01 ^a	4.33±0.23 ^b	1.03±0.05 ^b	4.43±0.23 ^b	1.44±0.10 ^c
DME	14.09±0.60 ^b	141.80±3.91 ^b	19.98±0.55 ^b	2.54±0.00 ^a	3.07±0.63 ^b	0.43±0.09 ^a	6.08±0.42 ^c	0.62±0.03 ^b
DAE	9.57±0.50 ^a	96.52±8.53 ^a	9.24±0.82 ^a	5.11±0.58 ^b	1.32±0.53 ^a	0.13±0.05 ^a	1.62±0.09 ^a	0.16±0.00 ^a

651 Data represent mean ± SD of triplicate analyses: different superscript letters (^{a–e}) with in a column indicate significant differences
 652 between the extracts ($P < 0.05$). Abbreviations: LL, lingonberry leaves; TPC, total phenolic content; WE, water extract; ME, methanol
 653 extract; AE, acetone extract; DME, deodorized methanol extract; DAE, deodorized acetone extract; edw, extract dry weight; pdw,
 654 plant dry weight.

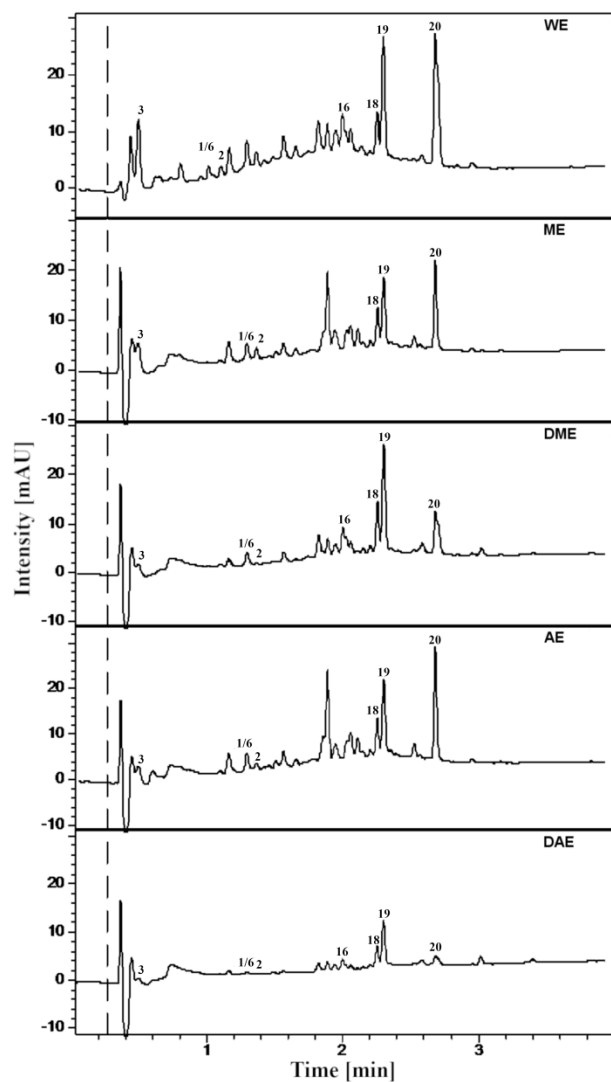
655

656 **Table 2– Compounds identified in the extracts of *Vaccinium vitis-idaea* L. leaves and their UPLC/qTOF–MS/MS data and the**
657 **content**

No. Compound	t _R (min)	[M-H] [–] (m/z)	Molecular ion formula	MS/MS (m/z)	Lingonberry leaves extracts				
					WE	ME	AE	DME	DAE
1 Catechin ¹	1.28	289.0719	C ₁₅ H ₁₃ O ₆	245.0826(100), 179.0306(25)	31.85±0.23 ^c	32.18±0.56 ^c	36.48±0.31 ^d	25.37±0.08 ^b	5.93±0.15 ^a
2 Epicatechin ²	1.55	289.0716	C ₁₅ H ₁₃ O ₆	245.0825(60), 179.0306(19)	14.62±0.35 ^a	32.18±0.57 ^b	14.33±0.16 ^a	18.56±9.5 ^{ab}	1.53±0.01 ^a
Total Flavan-3-ols					46.47	64.36	50.81	43.93	7.46
3 Quinic acid ¹	0.35	191.0564	C ₇ H ₁₁ O ₆	–	42.81±2.52 ^c	25.46±0.56 ^b	2.00±0.02 ^a	1.61±0.12 ^a	TR
4 Citric acid ¹	0.43	191.0201	C ₆ H ₇ O ₇	111.0083(20)	46.48±0.32	ND	ND	ND	ND
5 Caffeoyl-shikimic acid ²	1.08	335.0775	C ₂₀ H ₁₅ O ₅	179.0340(55), 161.0237(100)	2.85±0.02 ^e	2.22±0.10 ^d	1.96±0.02 ^c	1.25±0.02 ^b	0.72±0.05 ^a
6 Chlorogenic acid ¹	1.33	353.0874	C ₁₆ H ₁₇ O ₉	–	3.44±0.05 ^a	TR	TR	TR	ND
7 Caffeic acid ¹	1.47	179.0353	C ₉ H ₇ O ₄	–	2.62±0.04 ^b	ND	ND	1.04±0.00 ^a	0.98±0.08 ^a
8 <i>p</i> -Coumaric acid ¹	1.87	163.0402	C ₉ H ₇ O ₃	–	1.56±0.16 ^b	ND	ND	0.92±0.02 ^a	1.57±0.04 ^b
Total hydroxycinnamic acids					99.76	27.68	3.96	4.82	3.27
9 Arbutin ²	0.49	271.0824	C ₁₂ H ₁₅ O ₇	108.0218(100), 109.0275(8)	33.18±0.03 ^d	31.78±0.67 ^d	23.00±0.43 ^c	17.48±0.08 ^b	9.61±0.05 ^a
10 2- <i>O</i> -caffeoylarbutin ^{2,3}	1.70	433.1147	C ₂₁ H ₂₁ O ₁₀	323.0778 (5), 179.0350 (30), 161.0245 (100), 135.0449 (10)	24.11±1.01 ^c	29.85±0.35 ^d	34.34±0.24 ^e	21.92±0.07 ^b	10.21±0.23 ^a
11 Caffeoyl acetyl arbutin ³	2.56	475.1242	C ₂₃ H ₂₃ O ₁₁	179.0347(15), 161.0244(100)	8.51±0.14 ^b	8.82±0.00 ^b	10.97±0.22 ^c	10.49±0.31 ^c	6.05±0.08 ^a
Total Arbutin derivatives					65.8	70.45	68.31	49.89	25.87
12 B-type proanthocyanidin ²	1.34	577.1336	C ₃₀ H ₂₅ O ₁₂	407.0757(68), 289.0709(100), 245.0807(21), 161.0233(15), 125.0240(35)	8.35±0.35 ^a	TR	TR	TR	ND
13 A-type proanthocyanidin ^{2,3}	1.95	575.1195	C ₃₀ H ₂₃ O ₁₂	539.0998(65), 407.0791(48), 285.0387(100), 125.0245(58)	3.67±0.03 ^c	0.66±0.03 ^b	0.66±0.19 ^b	0.49±0.00 ^{ab}	0.27±0.07 ^a
Total proanthocyanidins					12.02	0.66	0.66	0.49	0.27
14 Rutin ¹	1.84	609.1459	C ₂₇ H ₂₉ O ₁₆	300.0267(100), 301.0332(30)	1.53±0.03 ^a	TR	TR	TR	ND
15 Quercetin-3- <i>O</i> -glucoside ¹	1.95	463.0881	C ₂₁ H ₁₉ O ₁₂	301.0339(66), 300.0272(100)	3.33±0.05 ^b	3.31±0.05 ^b	3.15±0.20 ^b	3.26±0.11 ^b	1.12±0.09 ^a

16	Quercetin pentoside ^{1,2,3}	2.10	433.0773	C ₂₀ H ₁₇ O ₁₁	301.0337(56), 300.0266(100), 271.0225(5)	20.43±0.07 ^c	ND	ND	18.20±0.32 ^b	7.70±0.25 ^a
17	Quercetin pentoside ^{2,3}	2.25	433.0769	C ₂₀ H ₁₇ O ₁₁	301.0345(58), 300.0274(100), 271.0238(5)	7.71±0.28 ^b	7.44±0.11 ^b	7.68±0.16 ^b	9.34±0.26 ^c	4.17±0.17 ^a
18	Quercetin-deoxyhexoside ^{2,3}	2.28	447.0931	C ₂₁ H ₁₉ O ₁₁	301.0338(23), 300.0252(100), 271.0227(86)	15.95±0.08 ^e	15.24±0.18 ^d	12.38±0.14 ^c	10.80±0.07 ^b	7.70±0.00 ^a
19	Quercetin-3- <i>O</i> -(HMG)- rhamnoside ³	2.67	591.1355	C ₂₇ H ₂₇ O ₁₅	529.1334(27), 489.1028(100), 447.0924(72), 301.0269(65)	29.80±0.10 ^d	0.41±0.03 ^b	0.61±0.02 ^c	0.29±0.01 ^a	0.30±0.01 ^a
20	Kaempferol-3- <i>O</i> -(HMG)- rhamnoside ³	2.95	575.1392	C ₂₇ H ₂₇ O ₁₄	515.1377(15), 473.1075(40), 431.0969(38), 285.0391(100)	1.78±0.02 ^d	1.07±0.02 ^b	1.46±0.08 ^c	1.02±0.06 ^b	0.38±0.01 ^a
Total flavonols						80.53	27.47	25.28	42.91	21.37
Total phenols						304.58	190.62	149.02	142.04	58.24

Data represent mean (AU×10⁻⁷/mg edw) ± SD of triplicate analyses: different superscript letters (^{a-e}) with in a row indicate significant differences between the extracts (*P* < 0.05). ¹Compounds identified by comparing with standard. ²Compounds identified tentatively by calculated molecular formula and fragmentation patterns using METLIN and/or HMDB databases. ³Compounds identified tentatively by calculated molecular formula and fragmentation patterns comparing to literature data (Ek et al., 2006; Liu et al., 2017). Abbreviations: WE, water extract; ME, methanol extract; AE, acetone extract; DME, deodorized methanol extract; DAE, deodorized acetone extract; edw, extract dry weight; NI, not identified; —, not performed; ND, not detected; TR, traces.



667

668 **Figure 1**—UPLC-UV chromatograms at 254 nm of lingonberry leaf extracts. Peak numbers
 669 correspond to the compounds listed in Table 2. Abbreviations: WE, water extract; ME, methanol
 670 extract; AE, acetone extract; DME, deodorized methanol extract; DAE, deodorized acetone
 671 extract.

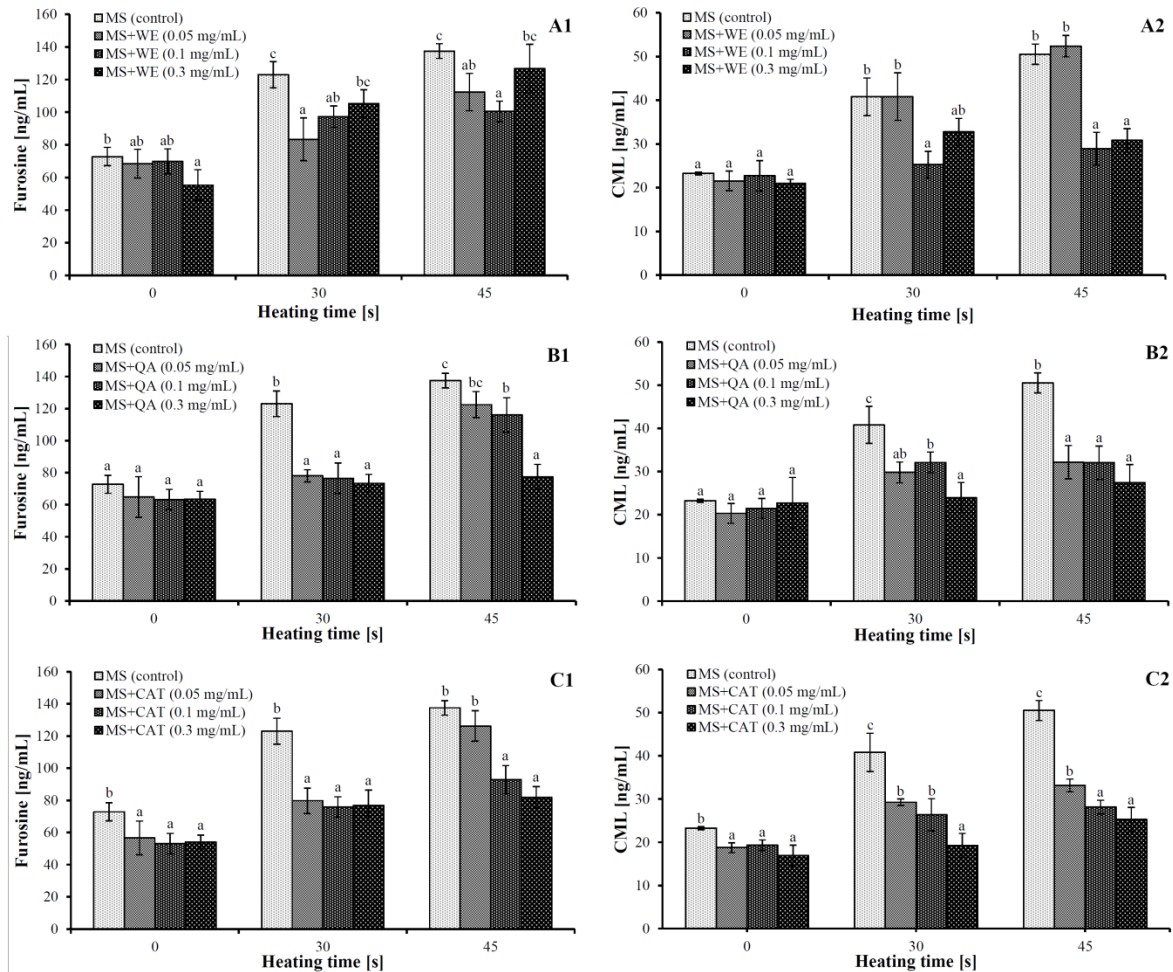


Figure 2—Effects of WE, quinic acid and catechin on furosine and CML formation in milk model system at different heating times: the different letters on the columns at the same heating time indicate the significant differences ($P < 0.05$). Abbreviations: MS, milk model system; QA, quinic acid; CAT, catechin; WE, water extract.

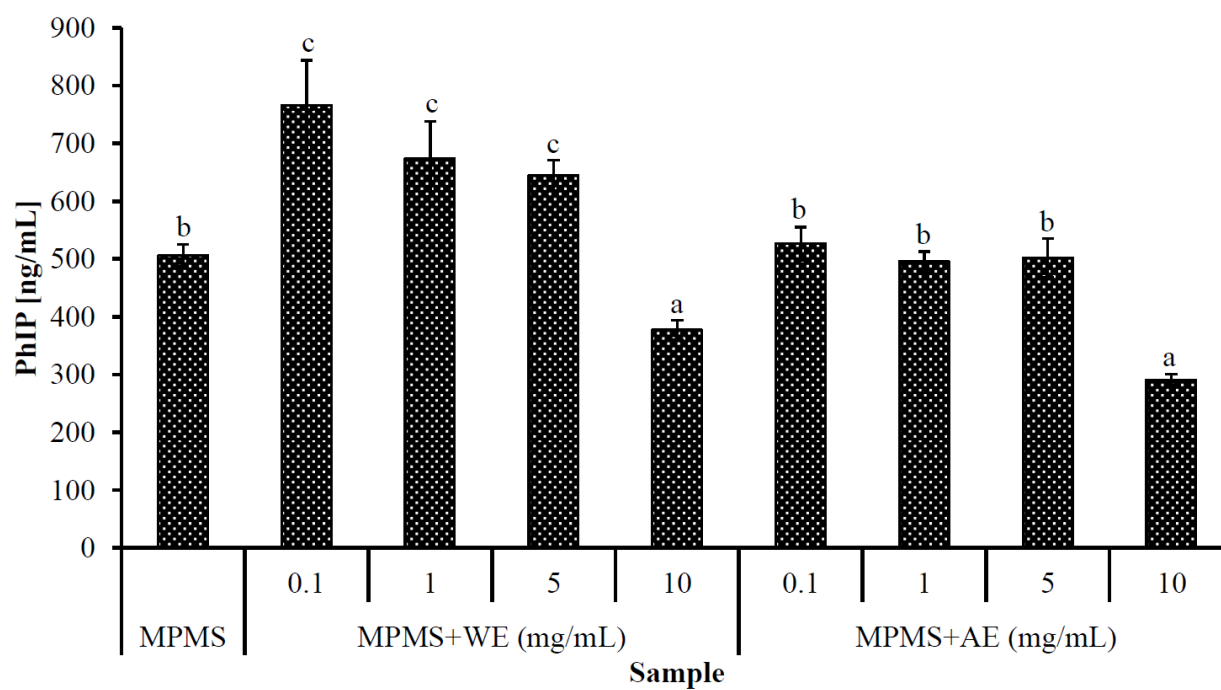


Figure 3—Effects of WE and AE of lingonberry leaves on PhIP formation in meat-protein model system. The different letters on the columns indicate the significant differences ($P < 0.05$). MPMS: meat-protein model system; WE: water extract; AE: acetone extract.

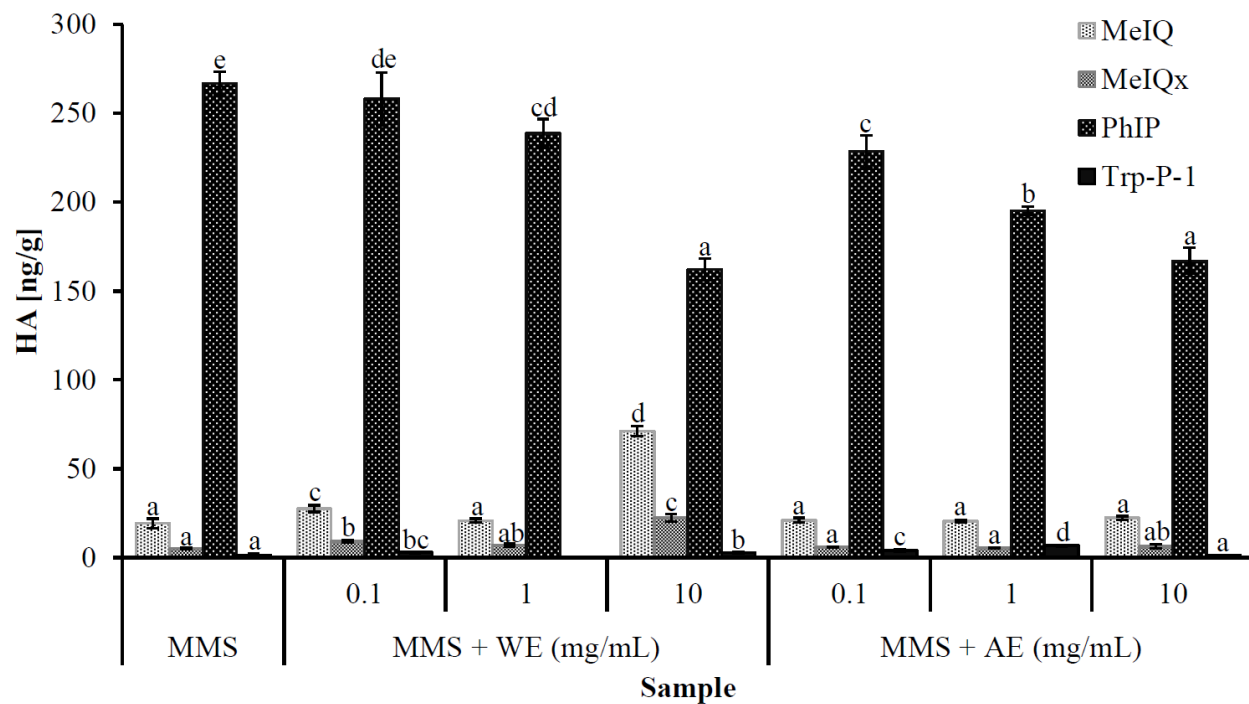


Figure 4—Effects of WE and AE of lingonberry leaves on HAs formation in meat model system. The different letters on the columns at the same HA indicate the significant differences ($P < 0.05$). MMS: meat model system; WE: water extract; AE: acetone extract.